



Mouse Dnmt3a DNA sequence

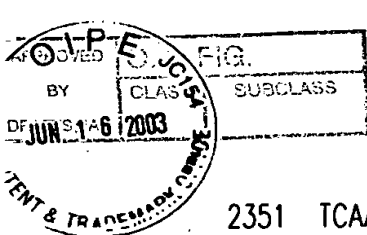
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 151 CTTCTCTGAA GCCCTCGCAG CCCACACAGG CCTCGCAGC CCCAGCCTGC
 201 CGCCTACTGC CCAGCAATGC CCTCCAGCGG CCCCGGGGAC ACCAGCAGCT
 251 CCTCTCTGGA GCGGGAGGAT GATCGAAAGG AAGGAGAGGA ACAGGAGGAG
 301 AACCGTGCCA AGGAAGAGCG CCAGGAGCCC AGCGCCACGG CCCGGAAGGT
 351 GGGGAGGCCT GGCCGGAAGC GCAAGCACCC ACCGGTGGA AGCAGTGACA
 401 CCCCCAAGGA CCCAGCAGTG ACCACCAAGT CTCAGCCCAT GGCCAGGAC
 451 TCTGGCCCCT CAGATCTGCT ACCCAATGGA GACTTGAGGA AGCGGAGTGA
 501 ACCCCAACCT GAGGAGGGGA GCCCAGCTGC AGGCCAGAAG GGTGGGGCCC
 551 CAGCTGAAGG AGAGGGAAGT GAGACCCAC CAGAAGCCTC CAGAGCTGTG
 601 GAGAATGGCT GCTGTGTGAC CAAGGAAGGC CGTGGAGCCT CTGCAGGAGA
 651 GGGCAAAGAA CAGAAGCAGA CCAACATCGA ATCCATGAAA ATGAGGGGCT
 701 CCCGGGGCCG ACTGCGAGGT GGCTTGGGCT GGGAGTCCAG CCTCCGTCAG
 751 CGACCCATGC CAAGACTCAC CTTCCAGGCA GGGGACCCCT ACTACATCAG
 801 CAAACGAAA CGGGATGAGT GGCTGGCAGC TTGAAAAGG GAGGCTGAGA
 851 AGAAAGCCAA GGTAATTGCA GTAATGAATG CTGTGGAAGA GAACCAGGCC
 901 TCTGGAGAGT CTCAGAAGT GGAGGAGGCC AGCCCTCCTG CTGTGCAGCA
 951 GCCCAGGAC CCTGCTTCTC CGACTGTGGC CACCACCCCT GAGCCAGTAG
 1001 GAGGGGATGC TGGGACAAG AATGCTACCA AAGCAGCCGA CGATGAGCCT
 1051 GAGTATGAGG ATGGCCGGGG CTTTGGCATT GGAGAGCTGG TGTGGGGGAA
 1101 ACTTCGGGGC TTCTCCTGGT GGCCAGGCCG AATTGTGTCT TGGTGGATGA

FIG. 1A-1



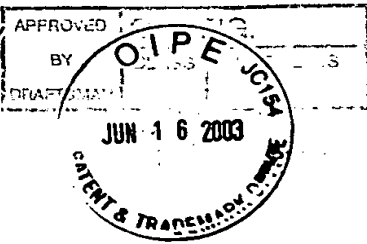
1151 CAGGCCGGAG CCGAGCAGCT GAAGGCACTC GCTGGGTCAT GTGGTTCGGA
1201 GATGGCAAGT TCTCAGTGGT GTGTGTGGAG AAGCTCATGC CGCTGAGCTC
1251 CTTCTGCAGT GCATTCCACC AGGCCACCTA CAACAAGCAG CCCATGTACC
1301 GCAAAGCCAT CTACGAAGTC CTCCAGGTGG CCAGCAGCCG TGCCGGAAG
1351 CTGTTTCCAG CTTGCCATGA CAGTGATGAA AGTGACAGTG GCAAGGCTGT
1401 GGAAGTGCAG AACAAGCAGA TGATTGAATG GGCCCTCGGT GGCTTCCAGC
1451 CCTCGGGTCC TAAGGGCCTG GAGCCACCAG AAGAAGAGAA GAATCCTTAC
1501 AAGGAAGTTT ACACCGACAT GTGGGTGGAG CCTGAAGCAG CTGCTTACGC
1551 CCCACCCCCA CCAGCCAAGA AACCAGAAA GAGCACAACA GAGAAACCTA
1601 AGGTCAAGGA GATCATTGAT GAGCGCACAA GGGAGCGGCT GGTGTATGAG
1651 GTGCGCCAGA AGTGCAGAAA CATCGAGGAC ATTTGTATCT CATGTGGGAG
1701 CCTCAATGTC ACCCTGGAGC ACCCACTCTT CATTGGAGGC ATGTGCCAGA
1751 ACTGTAAGAA CTGCTTCTTG GAGTGTGCTT ACCAGTATGA CGACGATGGG
1801 TACCAGTCCT ATTGCACCAT CTGCTGTGGG GGGCGTGAAG TGCTCATGTG
1851 TGGGAACAAC AACTGCTGCA GGTGCTTTTG TGTCCAGTGT GTGGATCTCT
1901 TGGTGGGGCC AGGAGCTGCT CAGGCAGCCA TTAAGGAAGA CCCCTGGAAC
1951 TGCTACATGT GCGGGCATAA GGGCACCTAT GGGCTGCTGC GAAGACGGGA
2001 AGACTGGCCT TCTCGACTCC AGATGTTCTT TGCCAATAAC CATGACCAGG
2051 AATTTGACCC CCCAAAGGTT TACCCACCTG TGCCAGCTGA GAAGAGGAAG
2101 CCCATCCGGG TGCTGTCTCT CTTTGATGGG ATTGCTACAG GGCTCCTGGT
2151 GCTGAAGGAC CTGGGCATCC AAGTGGACCG CTACATTGCC TCCGAGGTGT
2201 GTGAGGACTC CATCACGGTG GGCATGGTGC GGCACCAGGG AAAGATCATG
2251 TACGTCGGGG ACGTCCGCAG CGTCACACAG AAGCATATCC AGGAGTGGGG
2301 CCCATTGCAC CTGGTGATTG GAGGCAGTCC CTGCAATGAC CTCTCCATTG

FIG. 1A-2



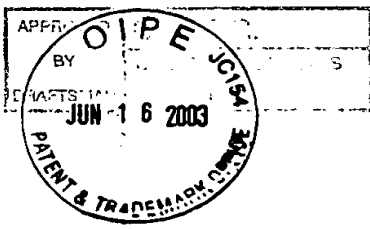
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2451 CCCCTTCTTC TGGCTCTTTG AGAATGTGGT GCCCATGGGC GTTAGTGACA
2501 AGAGGGACAT CTCGCGATTT CTTGAGTCTA ACCCCGTGAT GATTGACGCC
2551 AAAGAAGTGT CTGCTGCACA CAGGGCCCCG TACTTCTGGG GTAACCTTCC
2601 TGGCATGAAC AGGCCTTTGG CATCCACTGT GAATGATAAG CTGGAGCTGC
2651 AAGAGTGTCT GGAGCACGGC AGAATAGCCA AGTTCAGCAA AGTGAGGACC
2701 ATTACCACCA GGTCAAATC TATAAAGCAG GGCAAAGACC AGCATTTCCC
2751 CGTCTTCATG AACGAGAAGG AGGACATCCT GTGGTGCACT GAAATGGAAA
2801 GGGTGTTTGG CTTCCCCGTC CACTACACAG ACGTCTCCAA CATGAGCCGC
2851 TTGGCGAGGC AGAGACTGCT GGGCCGATCG TGGAGCGTGC CGGTCATCCG
2901 CCACCTCTTC GCTCCGCTGA AGGAATATTT TGCTTGTTG TAAGGGACAT
2951 GGGGGCAAAC TGAAGTAGTG ATGATAAAAA AGTTAAACAA ACAAACAAAC
3001 AAAAAACAAA ACAAACAAT AAAACACCAA GAACGAGAGG ACGGAGAAAA
3051 GTTCAGCACC CAGAAGAGAA AAAGGAATTT AAAGCAAACC ACAGAGGAGG
3101 AAAACGCCCG AGGGCTTGGC CTTGCAAAAG GGTGGACAT CATCTCCTGA
3151 GTTTTCAATG TTAACCTTCA GTCCTATCTA AAAAGCAAAA TAGGCCCTC
3201 CCCTTCTTCC CCTCCGTCC TAGGAGCGA ACTTTTGT TTTACTCTT
3251 TTTCAGAGG GTTTTCTGTT TGTTTGGTT TTTGTTTCTT GCTGTGACTG
3301 AAACAAGAGA GTTATTGCAG CAAAATCAGT AACAACAAAA AGTAGAAATG
3351 CCTGGAGAG GAAAGGGAGA GAGGGAAAAT TCTATAAAAA CTTAAATAT
3401 TGGTTTTTTT TTTTTTCT TTTCTATATA TCTCTTGGT TGTCTCTAGC
3451 CTGATCAGAT AGGAGCACAA ACAGGAAGAG AATAGAGACC CTCGGAGGCA
3501 GAGTCTCCTC TCCCACCCCC CGAGCAGTCT CAACAGCACC ATTCCTGGTC

FIG. 1A-3



3551 ATGCAAAACA GAACCCAAC AGCAGCAGGG CGCTGAGAGA ACACCACACC
3601 AGACACTTTC TACAGTATTT CAGGTGCCTA CCACACAGGA AACCTTGAAG
3651 AAAACCAGTT TCTAGAAGCC GCTGTTACCT CTTGTTTACA GTTTATATAT
3701 ATATGATAGA TATGAGATAT ATATATATAA AAGGTACTGT TAACTACTGT
3751 ACATCCCGAC TTCATAATGG TGCTTTCAAA ACAGCGAGAT GAGCAAAGAC
3801 ATCAGCTTCC GCCTGGCCCT CTGTGCAAAG GGTTTCAGCC CAGGATGGGG
3851 AGAGGGGAGC AGCTGGAGGG GGTTTAAACA AACTGAAGGA TGACCCATAT
3901 CACCCCCCAC CCCTGCCCCA TGCCTAGCTT CACCTGCCAA AAAGGGGCTC
3951 AGCTGAGGTG GTCGGACCCT GGGGAAGCTG AGTGTGGAAT TTATCCAGAC
4001 TCGCGTGCAA TAACCTTAGA ATATGAATCT AAAATGACTG CCTCAGAAAA
4051 ATGGCTTGAG AAAACATTGT CCCTGATTTT GAATTCGTCA GCCACGTTGA
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4151 CATTAACCCC NCCTGGAGCA AATAAAAAAA CATACAAAAT GT

FIG. 1A-4

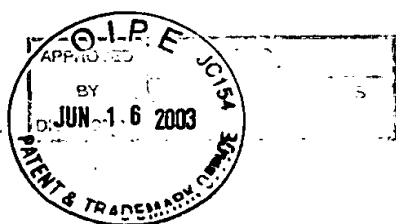


Mouse Dnmt3b1 DNA Sequence

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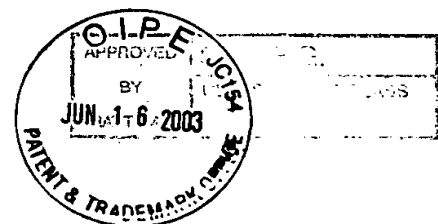
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101 ACCCGCGGCG CGATCGCGGC GCCGCGCTAC AGCCAGCCTC ACGACAGGCC
151 CGCTGAGGCT TGTGCCAGAC CTTGGAAACC TCAGGTATAT ACCTTTCCAG
201 ACGCGGGATC TCCCCTCCCC CATCCATAGT GCCTTGGGAC CAAATCCAGG
251 GCCTTCTTTC AGGAAACAAT GAAGGGAGAC AGCAGACATC TGAATGAAGA
301 AGAGGGTGCC AGCGGGTATG AGGAGTGCAT TATCGTTAAT GGGAACTTCA
351 GTGACCAGTC CTCAGACACG AAGGATGCTC CCTCACCCCC AGTCTTGGAG
401 GCAATCTGCA CAGAGCCAGT CTGCACACCA GAGACCAGAG GCCGCAGGTC
451 AAGCTCCCGG CTGTCTAAGA GGGAGGTCTC CAGCCTTCTG AATTACACGC
501 AGGACATGAC AGGAGATGGA GACAGAGATG ATGAAGTAGA TGATGGGAAT
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601 GACGCGCTCT GAAAGCCCGG CTGTCCGAAC CCGACATAGC AATGGGACCT
651 CCAGCTTGGA GAGGCAAAGA GCCTCCCCCA GAATCACCCG AGGTCCGCAG
701 GGCCGCCACC ATGTGCAGGA GTACCCTGTG GAGTTTCCGG CTACCAGGTC
751 TCGGAGACGT CGAGCATCGT CTTAGCAAG CACGCCATGG TCATCCCCTG
801 CCAGCGTCCA CTTATGGAA GAAGTGACAC CTAAGAGCGT CAGTACCCCA
851 TCAGTTGACT TGAGCCAGGA TGGAGATCAG GAGGGTATGG ATACCACACA
901 GGTGGATGCA GAGAGCAGAG ATGGAGACAG CACAGAGTAT CAGGATGATA
951 AAGAGTTTGG AATAGGTGAC CTCGTGTGGG GAAAGATCAA GGGCTTCTCC
1001 TGGTGGCCTG CCATGGTGGT GTCCTGAAA GCCACCTCCA ACCGACAGGC
  
```

FIG. 1B-1



1051 CATGCCCCGA ATGCGCTGGG TACAGTGGTT TGGTGATGGC AAGTTTTCTG
1101 AGATCTCTGC TGACAAACTG GTGGCTCTGG GGCTGTTTACG CCAGCACTTT
1151 AATCTGGCTA CCTTCAATAA GCTGGTTTCT TATAGGAAGG CCATGTACCA
1201 CACTCTGGAG AAAGCCAGGG TTCGAGCTGG CAAGACCTTC TCCAGCAGTC
1251 CTGGAGAGTC ACTGGAGGAC CAGCTGAAGC CCATGCTGGA GTGGGCCCCAC
1301 GGTGGCTTCA AGCCTACTGG GATCGAGGGC CTCAAACCCA ACAAGAAGCA
1351 ACCAGTGGTT AATAAGTCGA AGGTGCGTCG TTCAGACAGT AGGAACTTAG
1401 AAGCCAGGAG ACGCGAGAAC AAAAGTCGAA GACGCACAAC CAATGACTCT
1451 GCTGCTTCTG AGTCCCCCCC ACCCAAGCGC CTCAAGACAA ATAGCTATGG
1501 CGGGAAGGAC CGAGGGGAGG ATGAGGAGAG CCGAGAACGG ATGGCTTCTG
1551 AAGTCACCAA CAACAAGGGC AATCTGGAAG ACCGCTGTTT GTCCTGTGGA
1601 AAGAAGAACC CTGTGTCCTT CCACCCCCTC TTTGAGGGTG GGCTCTGTCA
1651 GAGTTGCCGG GATCGCTTCC TAGAGCTCTT CTACATGTAT GATGAGGACG
1701 GCTATCAGTC CTAAGTCACC GTGTGCTGTG AGGGCCGTGA ACTGCTGCTG
1751 TGCAGTAACA CAAGCTGCTG CAGATGCTTC TGTGTGGAGT GTCTGGAGGT
1801 GCTGGTGGGC GCAGGCACAG CTGAGGATGC CAAGCTGCAG GAACCCTGGA
1851 GCTGCTATAT GTGCCTCCCT CAGCGCTGCC ATGGGGTCCT CCGACGCAGG
1901 AAAGATTGGA ACATGCGCCT GCAAGACTTC TTCCTACTG ATCCTGACCT
1951 GGAAGAATTT GAGCCACCCA AGTTGTACCC AGCAATTCCT GCAGCCAAAA
2001 GGAGGCCCAT TAGAGTCCTG TCTCTGTTTG ATGGAATTGC AACGGGGTAC
2051 TTGGTGCTCA AGGAGTTGGG TATTAAAGTG GAAAAGTACA TTGCCTCCGA
2101 AGTCTGTGCA GAGTCCATCG CTGTGGGAAC TGTTAAGCAT GAAGGCCAGA
2151 TCAAATATGT CAATGACGTC CGGAAAATCA CCAAGAAAAA TATTGAAGAG
2201 TGGGGCCCGT TCGACTTGGT GATTGGTGGA AGCCCATGCA ATGATCTCTC

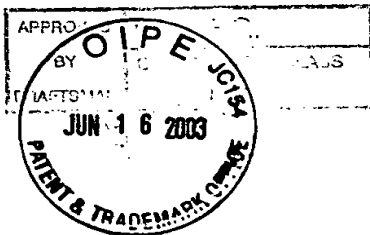
FIG. 1B-2



Appl. No. 09/720,086; 102(e): July 23, 2001
Dkt. No. 0609.4560002/JAG/KRM/DJN; Group Art Unit: 1642
Inventors: Li et al.; Tel: 202/371-2600
Title: *De Novo* DNA Cytosine Methyltransferase Genes, Polypeptides
and Uses Thereof

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2301 TCTTCGAGTT TTACCACTTG CTGAATTATA CCCGCCCCAA GGAGGGCGAC
2351 AACCGTCCAT TCTTCTGGAT GTTCGAGAAT GTTGTGGCCA TGAAAGTGAA
2401 TGACAAGAAA GACATCTCAA GATTCCTGGC ATGTAACCCA GTGATGATCG
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2551 GCTGCAGGAC TGCCTGGAGT TCAGTAGGAC AGCAAAGTTA AAGAAAGTGC
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3001 CTGGCCCTTG CAGGGGAGCC CCGGTGCTCC CTCCTGTGC ACAGCTCAGA
3051 CCTGGCTGCT TAGAGTAGCC CGGCATGGTG CTCATGTTCT CTTACCCTGA
3101 AACTTTAAAA CTTGAAGTAG GTAGTAAGAT GGCTTTCTTT TACCCTCCTG
3151 AGTTTATCAC TCAGAAGTGA TGGCTAAGAT ACCAAAAAAA CAAACAAAAA
3201 CAGAAACAAA AAACAAAAAA AAACCTCAAC AGCTCTCTTA GTACTCAGGT
3251 TCATGCTGCA AAATCACTTG AGATTTTGTT TTTAAGTAAC CCGTGCTCCA
3301 CATTTGCTGG AGGATGCTAT TGTGAATGTG GGCTCAGATG AGCAAGGTCA
3351 AGGGGCCAAA AAAAATTCCC CCTCTCCCCC CAGGAGTATT TGAAGATGAT
3401 GTTTATGGTT TAAGTCTTCC TGGCACCTTC CCCTTGCTTT GGTACAAGGG

FIG. 1B-3



Appl. No. 09/720,086; 102(e): July 23, 2001
 Dkt. No. 0609.4560002/JAG/KRM/DJN; Group Art Unit: 1642
 Inventors: Li et al.; Tel: 202/371-2600
 Title: *De Novo* DNA Cytosine Methyltransferase Genes, Polypeptides
 and Uses Thereof

3451 CTGAAGTCCT GTTGGTCTTG TAGCATTTC CAGGATGATG ATGTCAGCAG
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 3601 TGGAGTGGCC CTCTTCAGG TGTGAGGGAT ACGAAGGAGG AAGCTTAGGG
 3651 AAATCCATT C CCACTCCCT CTTGCCAAAT GAGGGGCCCA GTCCCAACA
 3701 GCTCAGGTCC CCAGAACCCC CTAGTTCCTC ATGAGAAGCT AGGACCAGAA
 3751 GCACATCGTT CCCCTTATCT GAGCAGTGT TGGGGAATA CAGTGAAAAC
 3801 CTTCTGGAGA TGTTAAAAGC TTTTACCCC ACGATAGATT GTGTTTTTAA
 3851 GGGGTGCTTT TTTTAGGGGC ATCACTGGAG ATAAGAAAGC TGCATTTTCA
 3901 AAATGCCATC GTAATGGTTT TTAACACCT TTTACCTAAT TACAGGTGCT
 3951 ATTTTATAGA AGCAGACAAC ACTTCTTTTT ATGACTCTCA GACTTCTATT
 4001 TTCA TGTTAC CATTTTTTTT GTA ACTCGCA AGGTGTGGGC TTTGTAACT
 4051 TCACAGGTGT GGGGAGAGAC TGCCTTGTTT CAACAGTTTG TCTCCACTGG
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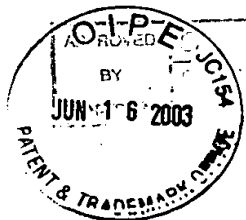
FIG. 1B-4



Human DNMT3A DNA Sequence

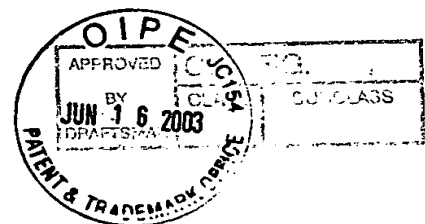
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28 CCGGCCCCGAC CCCACCGGCC ATACGGTGGG GCCATCGAAG CCCCCACCCA
78 CAGGCTGACA GAGGCACCGT TCACCAGAGG GCTCAACACC GGGATCTATG
128 TTTAAGTTTT AACTCTCGCC TCCAAAGACC ACGATAATTC CTTCCCCAAA
178 GCCCAGCAGC CCCCCAGCCC CGCGCAGCCC CAGCCTGCCT CCCGGCGCCC
228 AGATGCCCCG CATGCCCTCC AGCGGCCCCG GGGACACCAG CAGCTCTGCT
278 GCGGAGCGGG AGGAGGACCG AAAGGACGGA GAGGAGCAGG AGGAGCCGCG
328 TGGCAAGGAG GAGCGCCAAG AGCCCAGCAC CACGGCACGG AAGGTGGGGC
378 GGCCTGGGAG GAAGCGCAAG CACCCCCCGG TGGAAAGCGG TGACACGCCA
428 AAGGACCCTG CGGTGATCTC CAAGTCCCCA TCCATGGCCC AGGACTCAGG
478 CGCCTCAGAG CTATTACCCA ATGGGGACTT GGAGAAGCGG AGTGAGCCCC
528 AGCCAGAGGA GGGGAGCCCT GCTGGGGGGC AGAAGGGCGG GGCCCCAGCA
578 GAGGGAGAGG GTGCAGCTGA GACCCTGCCT GAAGCCTCAA GAGCAGTGA
628 AAATGGCTGC TGCACCCCCA AGGAGGGCCG AGGAGCCCTT GCAGAAGCGG
678 GCAAAGAACA GAAGGAGACC AACATCGAAT CCATGAAAAT GGAGGGCTCC
728 CGGGGCCGGC TCGGGGTGG CTTGGGCTGG GAGTCCAGCC TCGTCAGCG
778 GCCCATGCCG AGGCTCACCT TCCAGGCGGG GGACCCCTAC TACATCAGCA
828 AGCGCAAGCG GGACGAGTGG CTGGCAGCT GGAAAAGGA GGCTGAGAAG
878 AAAGCCAAGG TCACTGCAGG AATGAATGCT GTGGAAGAAA ACCAGGGGCC
928 CGGGGAGTCT CAGAAGGTGG AGGAGGCCAG CCCTCCTGCT GTGCAGCAGC
978 CCACTGACCC CGCATCCCCC ACTGTGGCTA CCACGCCTGA GCCCGTGGGG
1028 TCCGATGCTG GGGACAAGAA TGCCACCAA GCAGGCGATG ACGAGCCAGA

FIG. 1C-1



1078 GTACGAGGAC GGCCGGGGCT TTGGCATTGG GGAGCTGGTG TGGGGGAAAC
1128 TGGGGGGCTT CTCCTGGTGG CCAGGCCCGCA TTGTGTCTTG GTGGATGACG
1178 GGCCGGAGCC GAGCAGCTGA AGGCACCCGC TGGGTCATGT GGTTCGGAGA
1228 CGGCAAATTC TCAGTGGTGT GTGTTGAGAA GCTGATGCCG CTGAGCTCGT
1278 TTTGCAGTGC GTTCCACCAG GCCACGTACA ACAAGCAGCC CATGTACCGC
1328 AAAGCCATCT ACGAGGTCCT GCAGGTGGCC AGCAGCCGCG CGGGGAAGCT
1378 GTTCCCGGTG TGCCACGACA GCGATGAGAG TGACACTGCC AAGGCCGTGG
1428 AGGTGCAGAA CAAGCCCATG ATTGAATGGG CCCTGGGGGG CTTCACGCT
1478 TCTGGCCCTA AGGGCCTGGA GCCACCAGAA GAAGAGAAGA ATCCCTACAA
1528 AGAAGTGATC ACGGACATGT GGGTGAACC TGAGGCAGCT GCCTACGCAC
1578 CACCTCCACC AGCCAAAAG CCCCCGAAGA GCACAGCGGA GAAGCCCAAG
1628 GTCAAGGAGA TTATTGATGA GCGACAAGA GAGCGGCTGG TGTACGAGGT
1678 GCGGCAGAAG TGCCGGAACA TTGAGGACAT CTGCATCTCC TGTGGGAGCC
1728 TCAATGTTAC CCTGGAACAC CCCCTCTTCG TTGGAGGAAT GTGCCAAAAC
1778 TGCAAGAACT GCTTTCTGGA GTGTGCGTAC CAGTACGACG ACGACGGCTA
1828 CCAGTCCTAC TGCACCATCT GCTGTGGGGG CCGTGAGGTG CTCATGTGGC
1878 GAAACAACAA CTGCTGCAGG TGCTTTTGGC TGGAGTGTGT GGACCTCTTG
1928 GTGGGGCCGG GGGCTGCCCC GGCAGCCATT AAGGAAGACC CCTGGAAGTG
1978 CTACATGTGC GGGCACAAGG GTACCTACGG GCTGCTGCGG CGGCGAGAGG
2028 ACTGGCCCTC CCGGCTCCAG ATGTTCTTCG CTAATAACCA CGACCAGGAA
2078 TTTGACCCTC CAAAGGTTTA CCCACCTGTC CCAGCTGAGA AGAGGAAGCC
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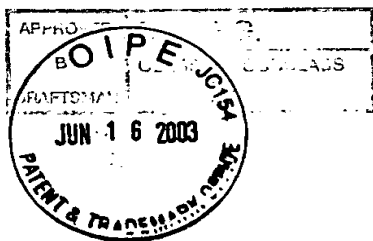
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2228 GAGGACTCCA TCACGGTGGG CATGGTGCGG CACCAGGGGA AGATCATGTA
2278 CGTCGGGGAC GTCCGCAGCG TCACACAGAA GCATATCCAG GAGTGGGGCC
2328 CATTGATCT GGTGATTGGG GGCAGTCCCT GCAATGACCT CTCCATCGTC
2378 AACCTGCTC GCAAGGGCCT CTACGAGGGC ACTGGCCGGC TCTTCTTTGA
2428 GTTCTACCGC CTCCTGCATG ATGCGCGGCC CAAGGAGGGA GATGATCGCC
2478 CCTTCTTCTG GCTCTTTGAG AATGTGGTGG CCATGGGCGT TAGTGACAAG
2528 AGGGACATCT CGCGATTTCT CGAGTCCAAC CCTGTGATGA TTGATGCCAA
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2628 GTATGAACAG GCCGTTGGCA TCCACTGTGA ATGATAAGCT GGAGCTGCAG
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2778 TCTTCATGAA TGAGAAAGAG GACATCTTAT GGTGCACTGA AATGGAAGG
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3078 CAGAAGAGAA AAAGGAATTT AAAACAAAAA CCACAGAGGC GGAAATACCG
3128 GAGGGCTTTG CCTTGCGAAA AGGGTTGGAC ATCATCTCCT GATTTTTCOA
3178 TGTTATTCTT CAGTCCTATT TAAAAACAAA ACCAAGCTCC CTTCCCTTCC
3228 TCCCCCTTCC CTTTTTTTTT GGTGAGACCT TTTATTTTCT ACTCTTTTCA
3278 GAGGGGTTTT CTGTTTGTGTT GGGTTTTGTT TCTGCTGTG ACTGAAACAA
3328 GAAGGTATT GCAGCAAAAA TCAGTAACAA AAAATAGTAA CAATACCTTG
3378 CAGAGGAAAG GTGGGAGGAG AGGAAAAAAG GGAAATTTTT AAAGAAATCT

FIG. 1C-3



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3428 ATATATTGGG TTGTTTTTTT TTTTGTITTT TGTTTTTTTT TTTTGGGTTT
 3478 TTTTTTTTTA CTATATAICT TTTTTTTGTT GTCTCTAGCC TGATCAGATA
 3528 GGAGCACAAG CAGGGGACGG AAAGAGAGAG AACTCAGGC GGCAGCATTC
 3578 CCTCCCAGCC ACTGAGCTGT CGTGCCAGCA CCATTCTGG TCACGCAAAA
 3628 CAGAACCCAG TTAGCAGCAG GGAGACGAGA ACACCACACA AGACATTTTT
 3678 CTACAGTATT TCAGGTGCCT ACCACACAGG AACCTTGAA GAAAATCAGT
 3728 TTCTAGAAGC CGCTGTTACC TCTGTTTAC AGTTTATATA TATATGATAG
 3778 ATATGAGATA TATATATAAA AGGTACTGTT AACTACTGTA CAACCCGACT
 3828 TCATAATGGT GCTTTCAAAC AGCGAGATGA GTAAAACAT CAGCTTCCAC
 3878 GTTGCCTTCT GCGCAAAGGG TTTCACCAAG GATGGAGAAA GGGAGACAGC
 3928 TTGCAGATGG CGCGTTCTCA CGGTGGGCTC TTCCCCTTGG TTTGTAACGA
 3978 AGTGAAGGAG GAGAACTTGG GAGCCAGGTT CTCCTGCCA AAAAGGGGGC
 4028 TAGATGAGGT GGTGGGGCCC GTGGACAGCT GAGAGTGGGA TTCATCCAGA
 4078 CTCATGCAAT AACCTTTGA TTGTTTTCTA AAAGGAGACT CCCTCGGCAA
 4128 GATGGCAGAG GGTACGGAGT CTTCAGGCCC AGTTTCTCAC TTTAGCCAAT
 4178 TCGAGGGCTC CTTGTGGTGG GATCAGAACT AATCCAGAGT GTGGGAAAGT
 4228 GACAGTCAAA ACCCCACCTG GAGCAAATAA AAAACATAC AAAACGTAAA
 4278 AAAAAAAAAA AAAAAA

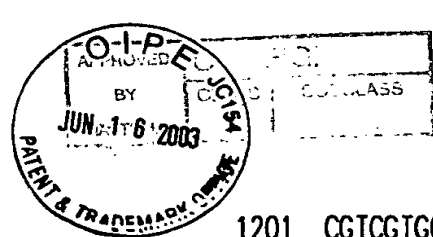
FIG. 1C-4



Human DNMT3B1 DNA Sequence:

1 GGCCGCGAAT TCGGCACGAG CCCTGCACGG CCGCCAGCCG GCCTCCCGCC
 51 AGCCAGCCCC GACCCGCGGC TCGCCGCCCC AGCCGCGCCC CAGCCAGCCC
 101 TCGGCAGGA AAGCATGAAG GGAGACACCA GGCATCTCAA TGGAGAGGAG
 151 GACGCCGGCG GGAGGAAGA CTCGATCCTC GTCAACGGGG CCTGCAGCGA
 201 CCAGTCCTCC GACTCGCCCC CAATCCTGGA GGCTATCCGC ACCCCGGAGA
 251 TCAGAGGCCG AAGATCAAGC TCGCGACTCT CCAAGAGGGA GGTGTCCAGT
 301 CTGCTAAGCT ACACACAGGA CTTGACAGGC GATGGCGACG GGAAGATGG
 351 GGATGGCTCT GACACCCAG TCATGCCAAA GCTCTTCGG GAAACCAGGA
 401 CTCGTTGAGA AAGCCCAGCT GTCCGAAGTC GAAATAACAA CAGTGTCTCC
 451 AGCCGGGAGA GGCACAGGCC TTCCCCACGT TCCACCCGAG GCCGGCAGGG
 501 CCGCAACCAT GTGGACGAGT CCCCCGTGGA GTTCCCGGCT ACCAGGTCCC
 551 TGAGACGGCG GGCAACAGCA TCGGCAGGAA CGCCATGGCC GTCCCCTCCC
 601 AGCTCTTACC TTACCATCGA CCTCACAGAC GACACAGAGG ACACACATGG
 651 GACGCCCCAG AGCAGCAGTA CCCCCTACGC CCGCCTAGCC CAGGACAGCC
 701 AGCAGGGGGG CATGGAGTCC CCGCAGGTGG AGGCAGACAG TGGAGATGGA
 751 GACAGTTCAG AGTATCAGGA TGGGAAGGAG TTTGGAATAG GGGACCTCGT
 801 GTGGGAAAG ATCAAGGGCT TCTCCTGGTG GCCCGCCATG GTGGTGTCTT
 851 GGAAGGCCAC CTCCAAGCGA CAGGCTATGT CTGGCATGCC GTGGGTCCAG
 901 TGGTTTGGCG ATGGCAAGTT CTCCGAGGTC TCTGCAGACA AACTGGTGGC
 951 ACTGGGGCTG TTCAGCCAGC ACTTTAATTT GGCCACCTTC AATAAGCTCG
 1001 TCTCCTATCG AAAAGCCATG TACCATGCTC TGGAGAAAGC TAGGGTGCGA
 1051 GCTGGCAAGA CTTCCCCAG CAGCCCTGGA GACTCATTGG AGGACCAGCT
 1101 GAAGCCCATG TTGGAGTGGG CCCACGGGGG CTTCAAGCCC ACTGGGATCG
 1151 AGGGCCTCAA ACCCAACAAC ACGCAACCAG TGGTTAATAA GTCGAAGGTG

FIG. 1D-1



1201 CGTCGTGCAG GCAGTAGGAA ATTAGAATCA AGGAAATACG AGAACAAGAC
 1251 TCGAAGACGC ACAGCTGACG ACTCAGCCAC CTCTGACTAC TCCCCCGCAC
 1301 CCAAGCGCCT CAAGACAAAT TGCTATAACA ACGGCAAAGA CCGAGGGGAT
 1351 GAAGATCAGA GCCGAGAACA AATGGCTTCA GATGTTGCCA ACAACAAGAG
 1401 CAGCCTGGAA GATGCCTGTT TGTCTTGTGG CAGGAAAAAC CCCGTGTCCT
 1451 TCCACCCTCT CTTTGAGGGG GGGCTCTGTC AGACATGCCG GGATCGCTTC
 1501 CTTGAGCTGT TTTACATGTA TGATGACGAT GGCTATCAGT CTTACTGCAC
 1551 TGTGTGCTGC GAGGCCCGAG AGCTGCTGCT TTGCAGCAAC ACGAGCTGCT
 1601 GCCGGTGTTT CTGTGTGGAG TGCCTGGAGG TGCTGGTGGG CACAGGCACA
 1651 GCGGCCGAGG CCAAGCTTCA GGAGCCCTGG AGCTGCTACA TGTGTCTCCC
 1701 GCAGCGCTGT CATGGCGTCC TGCGGCGCCG GAAGGACTGG AACGTGCGCC
 1751 TGCAGGCCTT CTTCAACAGT GACACGGGGC TTGAATACGA AGCCCCAAG
 1801 CTGTACCCTG CCATTCCCGC AGCCCGAAGG CGGCCCATTC GAGTCCTGTC
 1851 ATTGTTTGAT GGCATCGGA CAGGCTACCT AGTCTCAA A GAGTTGGCA
 1901 TAAAGGTAGG AAAGTACGTC GCTTCTGAAG TGTGTGAGGA GTCCATTGCT
 1951 GTTGAACCG TGAAGCACGA GGGGAATATC AAATACGTGA ACGACGTGAG
 2001 GAACATCACA AAGAAAAATA TTGAAGAATG GGGCCCATTT GACTTGGTGA
 2051 TTGGCGGAAG CCCATGCAAC GATCTCTCAA ATGTGAATCC AGCCAGGAAA
 2101 GGCCTGTATG AGGGTACAGG CCGGCTCTTC TTCGAATTTT ACCACCTGCT
 2151 GAATTACTCA CCCCCAAGG AGGGTGATGA CCGGCCGTTT TCTGGATGT
 2201 TTGAGAATGT TGTAGCCATG AAGGTTGGCG ACAAGAGGGA CATCTCACGG
 2251 TTCCTGGAGT GTAATCCAGT GATGATTGAT GCCATCAAAG TTTCTGCTGC
 2301 TCACAGGGCC CGATACTTCT GGGGCAACCT ACCCGGGATG AACAGGCCCC
 2351 TGATAGCATC AAAGAATGAT AACTCGAGC TGCAGGACTG CTTGAATAC
 2401 AATAGGATAG CCAAGTTAAA GAAAGTACAG ACAATAACCA CCAAGTCGAA

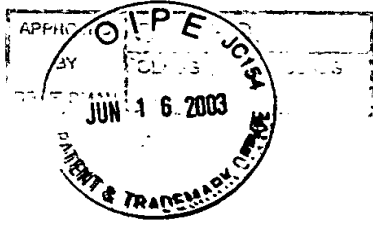
FIG. 1D-2



2451 CTCGATCAAA CAGGGGAAAA ACCAACTTTT CCCTGTTGTC ATGAATGGCA
2501 AAGAAGATGT TTTGTGGTGC ACTGAGCTCG AAAGGATCTT TGGCTTTCCT
2551 GTGCACTACA CAGACGTGTC CAACATGGGC CGTGGTGCCC GCCAGAAGCT
2601 GCTGGGAAGG TCCTGGAGCG TGCCTGTCAT CCGACACCTC TTCGCCCCCTC
2651 TGAAGGACTA CTTTGCATGT GAATAGTTCC AGCCAGGCCC CAAGCCCCT
2701 GGGGTGTGTG GCAGAGCCAG GACCCAGGAG GTGTGATTCC TGAAGGCATC
2751 CCCAGGCCCT GCTCTTCCTC AGCTGTGTGG GTCATACCGT GTACCTCAGT
2801 TCCCTCTTGC TCAGTGGGGG CAGAGCCACC TGA CTCTTGC AGGGGTAGCC
2851 TGAGGTGCCG CCTCCTTGTG CACAAATCAG ACCTGGCTGC TTGGAGCAGC
2901 CTAACACGGT GCTCATTTTT TCTTCTCCTA AAAC TT TAAA ACTTGAAGTA
2951 GGTAGCAACG TGGCTTTTTT TTTTCCCTT CCTGGGTCTA CCACTCAGAG
3001 AAACAATGGC TAAGATACCA AAACCACAGT GCCGACAGCT CTCCAATACT
3051 CAGGTTAATG CTGAAAATC ATCCAAGACA GTTATTGCAA GAGTTTAATT
3101 TTTGAAAAC TGGTACTGCT ATGTGTTTAC AGACGTGTGC AGTTGTAGGC
3151 ATGTAGCTAC AGGACATTTT TAAGGGCCCA GGATCGTTTT TTCCAGGGC
3201 AAGCAGAAGA GAAAATGTTG TATATGTCTT TTACCCGGCA CATTCCCCTT
3251 GCCTAAATAC AAGGGCTGGA GTCTGCACGG GACCTATTAG AGTATTTTCC
3301 ACAATGATGA TGATTTGAGC AGGGATGACG TCATCATCAC ATTCAGGGCT
3351 ATTTTTTCCC CCACAAACCC AAGGGCAGGG GCCACTCTTA GCTAAATCCC
3401 TCCCCGTGAC TGCAATAGAA CCCTCTGGGG AGCTCAGGAA GGGGTGTGCT
3451 GAGTTCTATA ATATAAGCTG CCATATATTT TGTAGACAAG TATGGCTCCT
3501 CCATATCTCC CTCTTCCCTA GGAGAGGAGT GTGAAGCAAG GAGCTTAGAT
3551 AAGACACCCC CTCAAACCCA TTCCCTCTCC AGGAGACCTA CCCTCCACAG
3601 GCACAGGTCC CCAGATGAGA AGTCTGCTAC CCTCATTTCT CATCTTTTTA
3651 CTAAACTCAG AGGCAGTGAC AGCAGTCAGG GACAGACATA CATTTCTCAT

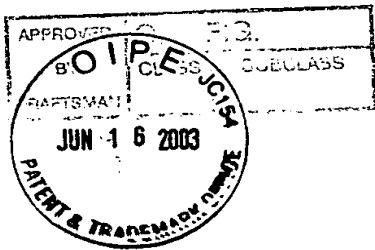
FIG. 1D-3

Appl. No. 09/720,086; 102(e): July 23, 2001
Dkt. No. 0609.4560002/JAG/KRM/DJN; Group Art Unit: 1642
Inventors: Li et al.; Tel: 202/371-2600
Title: *De Novo* DNA Cytosine Methyltransferase Genes, Polypeptides
and Uses Thereof



3701 ACCTTCCCCA CATCTGAGAG ATGACAGGGA AACTGCAAA GCTCGGTGCT
3751 CCCTTTGGAG ATTTTTTAAT CCTTTTTTAT TCCATAAGAA GTCGTTTTTA
3801 GGGAGAACGG GAATTCAGAC AAGCTGCATT TCAGAAATGC TGTCATAATG
3851 GTTTTAAACA CCTTTTACTC TTCTTACTGG TGCTATTTTG TAGAATAAGG
3901 AACAACTTG ACAAGTTTTG TGGGGCTTTT TATACACTTT TTAAAATCTC
3951 AACTTCTAT TTTTATGTTT AACGTTTCA TTAAAATTTT TTTGTAAGTG
4001 GAGCCACGAC GTAACAAATA TGGGGAAAAA ACTGTGCCTT GTTTCAACAG
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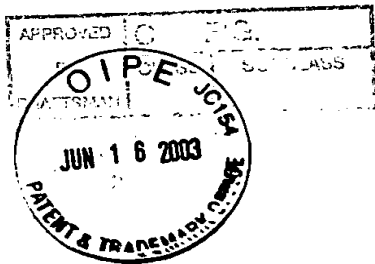
FIG. 1D-4



Mouse Dnmt3a Protein

1 MPSSGPGDTS SSSLEREDDR KEGEEQEENR GKEERQEPSA TARKVGRPGR
 51 KRKHPPVESS DTPKDAVTT KSQPMQDSG PSDLLPNGDL EKRSEPOPEE
 101 GSPAAGQKGG APAEGEGTET PPEASRAVEN GCCVTKEGRG ASAGEGKEQK
 151 QTNIESMKME GSRGRLRGGL GWESSLRQRP MPRLTFQAGD PYYISKRRKD
 201 EWLARWKREA EKKAKVIAVM NAVEENQASG ESQKVEEASP PAVQQPTDPA
 251 SPTVATTPEP VGGDAGDKNA TKAADDEPEY EDGRGFGIGE LWGKLRGFS
 301 WWPGRIVSWW MTGRSRAAEG TRWVMWFGDG KFSVVCVEKL MPLSSFCSAF
 351 HQATYNKQPM YRKAIYEVLQ VASSRAGKLF PACHDSDESD SGKAVEVQNK
 401 QMIEWALGGF QPSGPKGLEP PEECKNPYKE VYTDMMVEPE AAAYAPPPPA
 451 KKPRKSTTEK PKVKEIIDER TRERLVYEVN QKCRNIEDIC ISCGSLNVTL
 501 EHPLFIGGMC QNCKNCFLEC AYQYDDGYQ SYCTICCGGR EVLMCGNNNC
 551 CRCFCVECVD LLVGPGAAQA AIKEDPWNCY MCGHKGTYGL LRRREDWPSR
 601 LQMFFANNHD QEFDPPKVYP PVPAEKRKPI RVLSLFDGIA TGLLVKDLG
 651 IQVDRIASE VCEDSITVGM VRHQGKIMYV GDVRSVTQKH IQEWGPFDLV
 701 IGGSPCNDLS IVNPARKGLY EGTGRLEFFEF YRLLHDARPK EGDRPFFWL
 751 FENVVAMGVS DKRDISRFLE SNPVMIDAKE VSAHRARYF WGNLPGMNRP
 801 LASTVNDKLE LQECLEHGR I AKFSKVRTIT TRSNSIKQGK DQHFVFMNE
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 901 LKEYFACV*

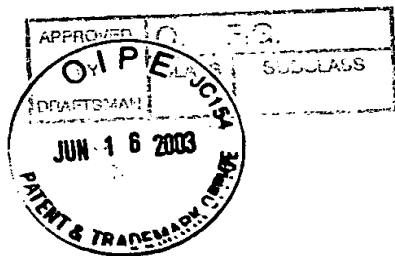
FIG. 2A



Mouse Dnmt3b1 Protein

1 MKGDSRHLNE EEGASGYEEC IIVNGNFSQ SSDTKDAPSP PVLEAICTEP
 51 VCTPETRGRR SSSRLSKREV SLLNYTQDM TGDGDRDDEV DDGNGSDILM
 101 PKLTRETKDT RTRSESPAVR TRHSNGTSSL ERQRASPRIT RGRQGRHHVQ
 151 EYPVEFPATR SRRRRASSSA STPWSSPASV DFMEEVTPKS VSTPSVDLSQ
 201 DGDQEGMDTT QVDAESRDGD STEYQDDKEF GIGDLVWGI KGFSSWWPAMV
 251 VSWKATSKRQ AMPGMRWWQW FGDGKFSEIS ADKLVALGLF SQHFNLATFN
 301 KLVSRYKAMY HTLEKARVRA GKTFSSSPGE SLEDQLKPLM EWAHGGFKPT
 351 GIEGLKPNKK QPVVNSKVR RSDSRNLEPR RRENKSRRRT TNSAASESP
 401 PPKRLKTNSY GKGDRGEDEE SRERMASEVT NNKGNLEDR LSCGKKNPVS
 451 FHPLFEGGLC QSCRDRFLEL FYMYDEGQY SYCTVCCEGR ELLLCNTSC
 501 CRFCFVECLE VLVGAGTAED AKLQEPWSCY MCLPQRCHGV LRRRKDWNMR
 551 LQDFFTTDPD LEEFEPKLY PAIPAAKRRP IRVLSLFDGI ATCYLVKEL
 601 GIKVEKYIAS EVCAESIAVG TVKHEGQIKY VNDVRKITKK NIEEWGPFDL
 651 VIGGSPCNDL SNVNPARKGL YEGTGRLEFE FYHLLNYTRP KEGDNRPFVW
 701 MFENVVAMKV NDKKDISRFL ACNPVMIDAI KVSAAHRARY FWGNLPGMNR
 751 PVMASKNDKL ELQDCLEFSR TAKLKKVQTI TTKSNSIRQG KNQLFPVVMN
 801 GKDDVLWCTE LERIFGFPAH YTDVSNMGRG ARQKLLGRSW SVPVIRHLFA
 851 PLKDYFACE*

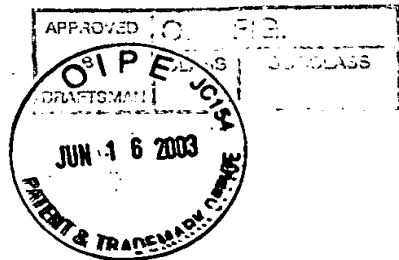
FIG. 2B



Human DNMT3A Protein

1 MPAMPSSGPG DTSSSAAERE EDRKDGEEQE EPRGKEERQE PSTTARKVGR
 51 PGRKRKHPPV ESGDTPKDPA VISKSPSMAQ DSGASELLPN GDLEKRSEPP
 101 PEEGSPAGGQ KGGAPAECEG AAETLPEASR AVENGCTPK EGRGAPAEAG
 151 KEQKETNIES MKMEGSRGRL RGGLGWESSL RQRMPRLTF QAGDPYYISK
 201 RKRDEWLARW KREAEEKKAV IAGMNAVEEN QGPGESQKVE EASPPAVQQP
 251 TDPASPTVAT TPEVGS DAG DKNATKAGDD EPEYEDGRGF GIGELVWGKL
 301 RGFSSWWPGRI VSWMTGRSR AAETRWWMW FGDGKFSVVC VEKLMPLSSF
 351 CSAFHQATYN KQPMYRKAIY EVLQVASSRA GKLFVCHDS DESDTAKAVE
 401 VQNKPMIEWA LGGFQPSGPK GLEPPEEEKN PYKEVYTDWW VEPEAAAYAP
 451 PPPAKKPRKS TAEKPKVKEI IDERTRERLV YEVRQKCRNI EDICISCGSL
 501 NVTLEHPLFV GGMCQNCKNC FLECAQYDD DGYQSYCTIC CGGREVLMCG
 551 NNNCCRCFCV ECVDLLVGPG AAQAAIKEDP WNCYMCCHKG TYGLLRRED
 601 WPSRLQMFFA NNHDQEFDP KVPVPVPAEK RKPIRVLSLF DGIATGLLV
 651 KDLGIQVDY IASEVCDSI TVGMVRHQGK IMYVGDVRSV TQKHIQEWGP
 701 FDLVIGGSPC NDLSIVNPAR KGLYEGTGR FFEFYRLLHD ARPKEGDDRP
 751 FFWLFENVVA MGVSDKRDIS RFLESNPVMI DAKEVSAHR ARYFWGNLPG
 801 MNRPLASTVN DKLELQECLE HGRIAKFSKV RTITTRSNSI KQKQDQHPV
 851 FMNEKEDILW CTEMERVFGF PVHYTDVSNM SRLARQRLG RSWSPVIRH
 901 LFAPLKEYFA CV*

FIG. 2C

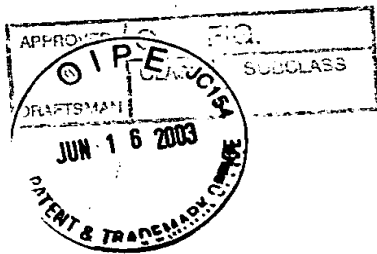


Appl. No. 09/720,086; 102(e): July 23, 2001
 Dkt. No. 0609.4560002/JAG/KRM/DJN; Group Art Unit: 1642
 Inventors: Li et al.; Tel: 202/371-2600
 Title: *De Novo* DNA Cytosine Methyltransferase Genes, Polypeptides
 and Uses Thereof

Human DNMT3B1 Protein

1 MKGDTRHLNG EEDAGGREDS ILVNGACSDQ SSDSPPILEA IRTPEIRGRR
 51 SSSRLSKREV SSLLSYTQDL TGDGDGEDGD GSDTPVMPKL FRETRTRSES
 101 PAVTRRNNS VSSRERHRPS PRSTRGRQGR NHVDESPVEF PATRSLRRRA
 151 TASAGTPWPS PPSSYLTLDL TDDTEDHGT PQSSSTPYAR LAQDSQQGGM
 201 ESPQVEADSG DGDSEYQDG KEFGIGDLVW GKIKGFSWMP AMVSWKATS
 251 KRQAMSGMRW VQWFGDGKFS EVSADKLVAL GLFSQHFNLA TFNKLVSRYK
 301 AMYHALEKAR VRAGKTFPSS PGDSLEDQLK PMLEWAHGGF KPTGIEGLKP
 351 NNTQPVVNKS KVRAGSRKL ESRKYENKTR RRTADDSATS DYCPAPKRLK
 401 TNCYNNGKDR GDEDQSREQM ASDVANKSS LEDGCLSCGR KNPVSFHPLF
 451 EGGLCQTCRD RFLELFMYD DDGYQSYCTV CCEGRELLLC SNTSCCRCFC
 501 VECLEVLVGT GTAAEAKLQE PWSCYMCLPQ RCHGVLRRRK DWNVRLQAFF
 551 TSDTGLEYES PKLYPAIPAA RRRPIRVLSL FDGIATGYLV LKELGIKVGK
 601 YVASEVCEES IAVGTVKHEG NIKYVNDVRN ITKKNIEEWG PFDLVIGGSP
 651 CNDLSNVNPA RKGLYEGTGR LFFEFYHLLN YSRPKEGDDR PFFWMFENVV
 701 AMKVGDKRDI SRFLECNPMV IDAIKVSAAH RARYFWGNLP GMNRPVIASK
 751 NDKLELQDCL EYNRIAKLKK VQTITTKSNS IKQGKNQLFP VMNGKEDVL
 801 WCTELERIFG FPVHYTDVSN MGRGARQKLL GRWSVPVIR HLFAPLKDYF
 851 ACE*

FIG. 2D

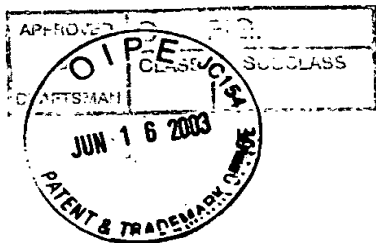


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 Dkt. No. 0609.4560002/JAG/KRM/DJN; Group Art Unit: 1642
 Inventors: Li et al.; Tel: 202/371-2600
 Title: *De Novo* DNA Cytosine Methyltransferase Genes, Polypeptides
 and Uses Thereof

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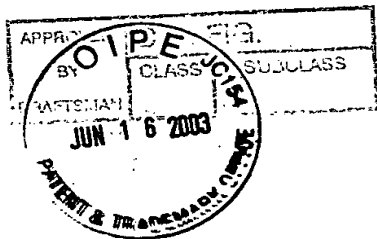
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Dnmt3a  51  KRKHPPVSSDTPKDPVTTKSQPMADSGPSD....LLPNGDLEKRSEP  96
      |. : : | | : : | |
Dnmt3b  1  .....MKGDSRHLNEEGASGYEECIIVNGNFSDDQSSD  33
Dnmt3a  97  QPEEGSP....AAGQKGGAPAEGETETPPEAS.RAVENGCCVTKE..GR  139
      : || | | | | | | | | | | | | | | | |
Dnmt3b  34  TKDAPSPPVLEAICTEPVCTPETRGRSSRLSKREVSSLLNYTQDMTGD  83
Dnmt3a  140 G.....ASAGEG.....KEQKQTNIESMKMEGSRGRLRGGLGWESSLRQ  178
      | | | | | | | | | | | | | | | | | |
Dnmt3b  84  GDRDDEVDDGNGSDILMPKLTREKDTTRSESPAVRTRHSNGTSSLERQ  133
Dnmt3a  179 RMPRLTFQAGDPYYISKRKDEWLARWKREAEEKKAKVIAVMNAVEENQA  228
      | ||:| : : | : . : : . . |
Dnmt3b  134 RASPRITRGRQGRHHV.....QEYPVEFPATRSRRRRASSASTPWSSPA  178
Dnmt3a  229 SGESQKVEEASPPAVQQTDPASPTVATTPEPVGGDAGDKNATKAADDEP  278
      | : . || . | . | | | | | | | | | |
Dnmt3b  179 SVDF..MEEVTPKSVSTP....SVDLSQGDQEGMDTTQVDAESRDGDST  222
Dnmt3a  279 EYEDGRGFGIGELVWGKLRGFSWWPGRIVSWWMTGRSRAAEGTRWVMWFG  328
      ||:| : ||||:|||||:||||| :||| | : . | ||| |||
Dnmt3b  223 EYQDDKEFGIGDLVWGKIKGFSWWPAMVVSWKATSKRQAMPGRWVQWFG  272
Dnmt3a  329 DGKFSVVCVEKLMLPLSSFCSAFHQATYNKQPMYRKAIYEV LQVASSRAGK  378
      ||||| : :||. | | | | | | | | | | | | | |
Dnmt3b  273 DGKFSEISADKLVALGLFSQHFNLATFNKLVSYRKAMYHTLEKARVRAGK  322
Dnmt3a  379 LFPACHDSDES DSGKAVEVQNKQMI EWALGGFQPSGPKGLEPPEEEK..N  426
      | | | | | | | | | | | | | | | | | | | |
Dnmt3b  323 TF.....SSSPGESLEDQLKPMLEWAHGGFKPTGIEGLKPNKKQPVVN  365
Dnmt3a  427 PYKEVYTD MW.VEP.....EAAAYAPPPPAKKPRKSTTEKPK  462
      | . | . || : || | | | : | | |
Dnmt3b  366 KSKVRRSDSRNLEPRRRENKSRRTTND SAASESPPPKRLKTNSYGGKDR  415
  
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FIG.3A-1



Dnmt3a	463	VKEIIDERTRERLVYEVROKCRNI	IEDICISCGSLNVTLEHPFFIGGMCQN	512
		. . :	: : :	
Dnmt3b	416	GE...DEESRERMASEVTNNKG	NLEDRLSCGKKNPVSFHPLFEGGLCQS	462
Dnmt3a	513	CKNCFLECAYYQDDGYQSYCTICCGGREVLMCGNNCCRCFCVECVDLL		562
		: . : : . : . . :		
Dnmt3b	463	CRDRFLELFYMYDEGYSYCTVCCEGRELLLCSTSCRCFCVECLEVL		512
Dnmt3a	563	VGPGAAQAAIKEDPWNCYMC	GHKGTGYLLRRREDWPSRLQMFFANNHD.Q	611
		: : . . : . . . :		
Dnmt3b	513	VGAGTAEDAKLQEPWSCYMC	LQRCHGVLRRRKDOWNMRLQDFFTTDPDLE	562
Dnmt3a	612	EFDPPKVYPVPAEKRP	IRVLSLFDGIATGLLVLKDLGIQVDRIASEV	661
		: . : : : . : :		
Dnmt3b	563	EFEPKLYPAIPAARRP	IRVLSLFDGIATGYLVKELGIKVEKYIASEV	612
Dnmt3a	662	CEDSITVGMVRHQKIMYVGDVRSVTQKH	IQEWGPFDLVIGGSPCNDLSI	711
		: : : . : . : :		
Dnmt3b	613	CAESIAVGTVKHEGQIKYVNDVRKITKKNIEEWGPFDLVIGGSPCNDLSN		662
Dnmt3a	712	VNPARKGLYEGTGRLFFEFYRLLHDARPKEGDDRPFFWLFENVVAMGVSD		761
		. . : .		
Dnmt3b	663	VNPARKGLYEGTGRLFFEFYHLLNYTRPKEGDNRPFFWMFENVVAMKVND		712
Dnmt3a	762	KRDISRFLESNPVMIDAKEVSAHRARYFWGNLPGMNRPLASTVNDKLEL		811
		: . . .		
Dnmt3b	713	KKDISRFLACNPVMIDAIVSAHRARYFWGNLPGMNRPMASKNDKLEL		762
Dnmt3a	812	QECLEHGRIAKFSKVRTITTRSNSIKQKGDQHF	PVFMNEKEDILWCTEME	861
		: . : : . : : :		
Dnmt3b	763	QDCLEFSRTAKLKKVQTITTKSNSIRQGNQLFPVVMNGKDDVLWCTELE		812
Dnmt3a	862	RVFGFPVHYTDVSNMSRLARQRLGRSWSVPVIRHLFAPLKEYFACV*		909
		: : :		
Dnmt3b	813	RIFGFPAHYTDVSNMGRGARQKLLGRSWSVPVIRHLFAPLKDYFACE*		860

FIG.3A-2



DNMT3A 1 MPAMPSSGPGDTSSSAAEREEDRKDGEEQEEPRGKEERQEPSTTARKVGR

DNMT3A 51 PGRKRKHPPVESGDTPKDPAVISKSPSMAQDSGASELLPNGDLEKRSEPO
 DNMT3B 1MKGDTRHLNGEEDAGGREDSILVNGACSDQSSDSP

DNMT3A 101 PEEGSPAGGQKGGAPAEEGEGAAETLPEASRAVENGCCTPKEGRGAPAEAG
 DNMT3B 36 PILEAIRTPAIRGGWASSRLSKREVSSLLSYTQDLTGDDGEDGDSOTP

DNMT3A 151 KEQKETNIESMKMEGSRGRLRGGLGWESSLRQRMPRLTFQAGDPYYISK
 DNMT3B 86 VMPKLFRETRTRSESPAVRTRNNNSVSSRERHRPSRSTRGRQGRNHVDE

DNMT3A 201 RKRDEWLARWKREAEEKAKVIAGMNAVEENQGPGESQKVEEASPPAVQQP
 DNMT3B 136 SPVEFPATRSLRRRATASAGTPWPSPSSYLTIDLTDDTEDTH..GTPQS

DNMT3A 251 TDPASPTVATTPEPVGSDAGDKNATKAGDDEPEYEDGRGFGIGELVWGKL
 DNMT3B 184 SSTPYARLAQDSQQGGMESPVQVEADSGDGDSEYQDGKEFGIGDLVWGKI

DNMT3A 301 RGFSWWPGRIVSWWMTGRSRAAEGTRWVMWFGDGKFSVVCVEKLMPLSSF
 DNMT3B 234 KGFSWWPAMVVSWKATSKRQAMSGMRWVQWFGDGKFSEVSADKLVALGLF

DNMT3A 351 CSAFHQATYNKQPMYRKAIYEVLQVASSRAGKLFVCHDSDESDTAKAVE
 DNMT3B 284 SQHFNLATFNKLVSRYKAMYHALEKARVRAGKTFP.....SSPGDSLE

DNMT3A 401 VQNKPMIEWALGGFQPSGPKGLEP....PEEEKNPYKEVYTDMWVE....
 DNMT3B 327 DQLKPMLEWAHGGFKPTGIEGLKPNNTQPVVNKSQVRRAGSRKLESRYE

DNMT3A 443PEAAAYAPPPAKKPRKSTAEKPKVKEIIDERTRERLVYEVRO
 DNMT3B 377 NKTRRRRTADSATS DYCPAPKRLKTNCYNNGKDRGDEDQSREQMASDVAN

FIG.3B-1

FIG. 3B-2

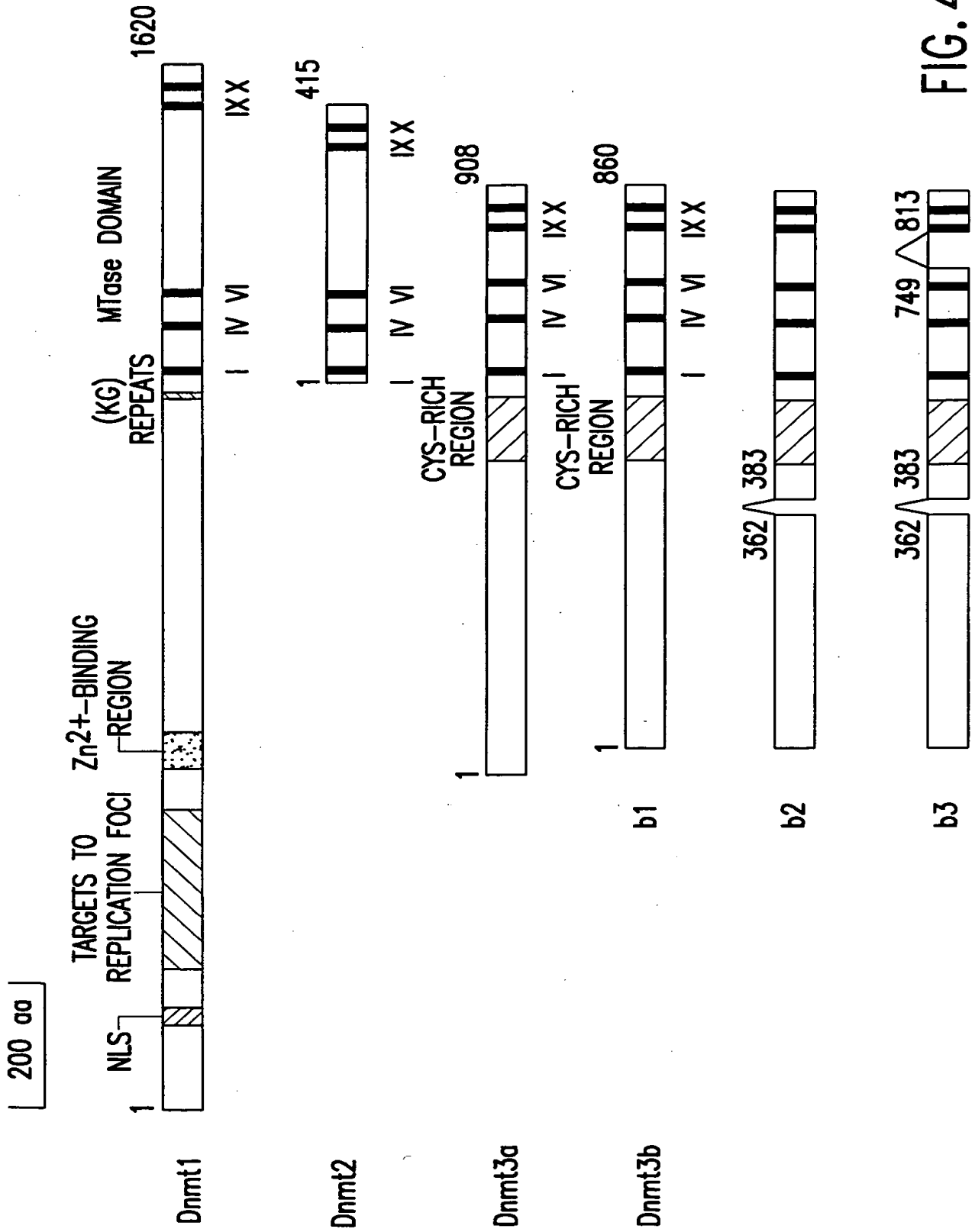
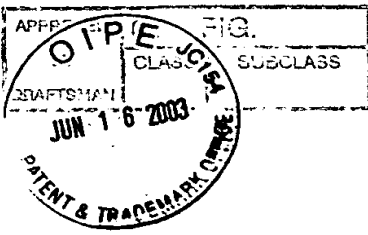


FIG. 4A

APPROVED	FIG.
BY	SUBCLASS



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 Dkt. No. 0609.4560002/JAG/KRM/DJN; Group Art Unit: 1642
 Inventors: Li et al.; Tel: 202/371-2600
 Title: *De Novo* DNA Cytosine Methyltransferase Genes, Polypeptides
 and Uses Thereof

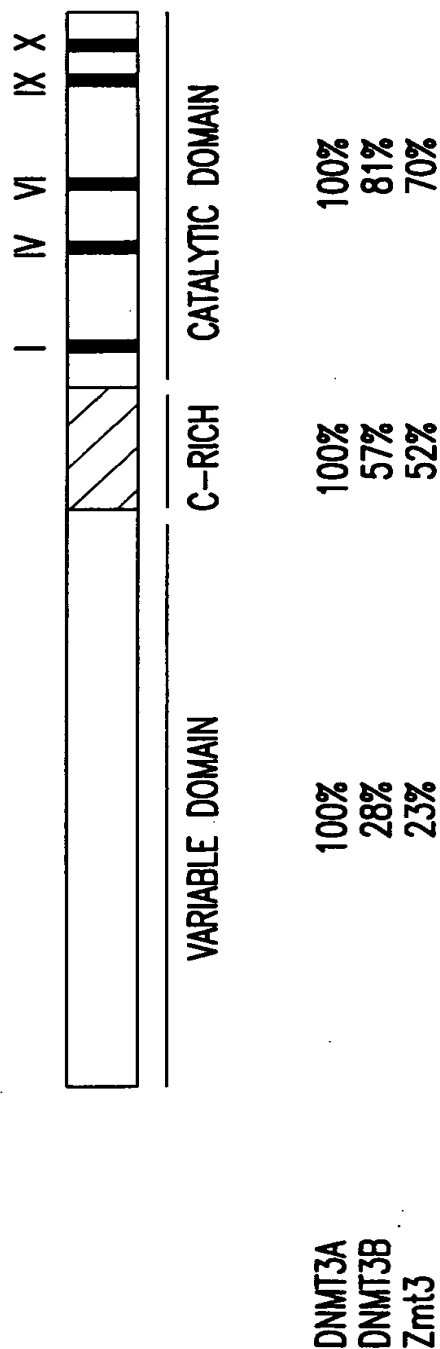


FIG. 4B

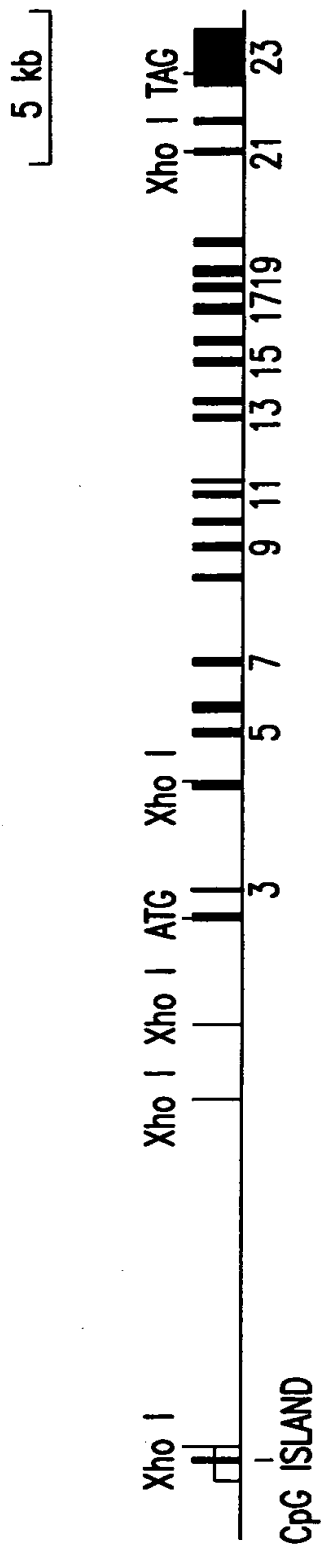
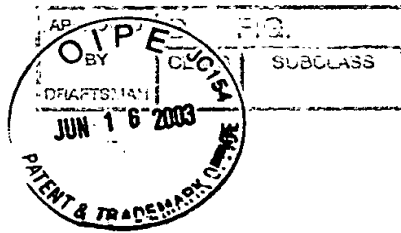
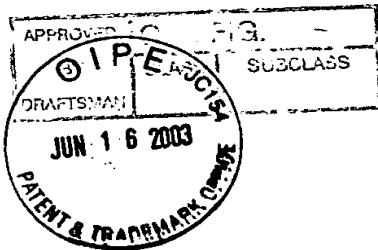


FIG. 4C

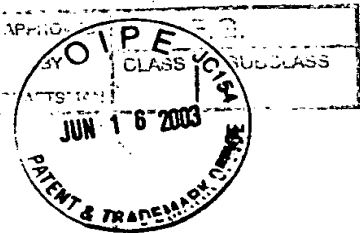


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Exon1 (>=90bp) CGGAGgtgagcgccccggg.intron(17618bp). tggcttctccacagGAAAGC
 Exon2 (148bp) TCAGAGgtgctgggcagtg.intron(887bp). CTGTTTCTCTACAGGCCGAA
 Exon3 (62bp) ACACAGgtatggtctctgtc.intron(3343bp). tgttccctataaagGACTTG
 Exon4 (102bp) CCAGCTgtaagtagccacacc.intron(1642bp). ctctcttgcttcttagGTCCGA
 Exon5 (125bp) ACCAGGgttggtcccccagatg.intron(602bp). tccctctgtccacagTCCCTG
 Exon6 (222bp) TATCAGgtatggccgagaggg.intron(1403bp). tgggttttcttccagGATGGG
 Exon7 (159bp) TCCGAGgtgagtcggggaag.intron(2588bp). gtcttctctttagGTCTCT
 Exon8 (108bp) CTGGAGgttaacatgggatgag.intron(917bp). actctgcctttgcagAAAGCT
 Exon9 (145bp) AACCAGgtgggaatgagtccc.intron(765bp). tttccctcctcaaaagTGGTTA
 Exon10 (60bp) AATACGgtatttctctctgt.intron(1813bp). aattaccttccacagAGAAACA
 Exon11 (126bp) GCCGAGgtgattgttgggtac.intron(115bp). ttcttttctcaatagAACAA
 Exon12 (45bp) TGGAGgttaacgttctctccc.intron(1095bp). ctgttttcttaccagATGGCT
 Exon13 (80bp) TGCCGGgttaagtctctctact.intron(417bp). ctctctggctgccagGATCGC
 Exon14 (113bp) CTGCCGgtgagcactgggccc.intron(1160bp). tgccactgggtccagGTGTTT
 Exon15 (184bp) GAATACgttaagccacaggtc.intron(600bp). ttccctacctggcagGAAGCC
 Exon16 (85bp) CGACAGgtgagttcggggaac.intron(824bp). ctctggccccccacagGCTACC
 Exon17 (146bp) AAAATgtgagggcagctctgt.intron(536bp). gtctctctcttccagATTGAA
 Exon18 (91bp) TGTATGgtgagcactctctc.intron(352bp). cttttctgagcacagAGGGTA
 Exon19 (149bp) CTGCAGgtgagggaaatctggg.intron(958bp). tctttctccccacagTGTAAAT
 Exon20 (86bp) GAACAGgttaacaaagggtct.intron(2867bp). tttggctgttccccagGCCCGT
 Exon21 (70bp) GCCAAGgttaagaaagtacag.intron(801bp). cattttgttctccagTTAAAG
 Exon22 (119bp) CGAAAGgtgagcaaggctgca.intron(1434bp). ctccgggtacccccagGATCTT
 Exon23 (1585bp)

FIG.4D

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	I	IV	VI
DNMT1	DVFSGCGGLSEGFHQAG	DVEMLCGGPPCQGFSGNMR	YRPRFFLLENVRNFVSFKR
Dnmt1	DVFSGCGGLSEGFHQAG	DVEMLCGGPPCQGFSGNMR	YRPRFFLLENVRNFVSFRR
MET1(Ath)	DIFAGCGGLSHGLKKAG	QVDFINGGPPCQGFSGNMR	FRPRYFLLLENVRTFVSFNK
Masc1	DTFCGGGVSLGARQAG	HVDILHSPPCQTFSSRAHT	VRPRLFTVEETDGIIMDRQS
Masc2	DIFAGCGGLTLGLDLSG	EVDFTYGGPPCQGFSGNMR	YKPRFVLLLENVKGLITTKL
Dnmt2	ELYSIGGMHHAALRESH	SFNMLMSPPCQPFTRIGL	KLPKYILLLENVKGFVSST
M.Spr	SLFSGIGAFEAALRNIG	EDLLVGGSPCQSFVAGH	KQPKFFVFENVKGLINHDK
DNMT3A	SLFDGIATGLLVKDLG	PFDLVIGGSPCNDLSIVNP	DRPFFWLLENVAMGVSDK
Dnmt3a	SLFDGIATGLLVKDLG	PFDLVIGGSPCNDLSIVNP	DRPFFWLLENVAMGVSDK
DNMT3B	SLFDGIATGYLVVKELG	PFDLVIGGSPCNDLSIVNP	DRPFFWLFENVAMKVGDK
Dnmt3b	SLFDGIATGYLVVKELG	PFDLVIGGSPCNDLSIVNP	NRPFWMFENVAMKVNDK
Zmt3	SLFDGIATGYLVLRDLG	PFDLLIGGSPCNDLSIVNP	POPFFWLFENUTFMQTHVK
consensus	--F-G-----G	---GG-PC---S-N	---P-F---ENW----

	IX	X
DNMT1	RVVSVRECARSCGFP	LFGNILDKHRQVGNVPPPLAKAIG
Dnmt1	RVVSVRECARSCGFP	FFGNILDRHRQVGNVPPPLAKAIG
MET1(Ath)	RILTVRECARSCGFP	FAGNINHKKRQIGNAVPPPLAFALG
Masc1	RKFTVRELACIQGFP	FVGTLTKRRIIGNAVPPPLSAAIM
Masc2	RVYTVRELARACGFP	GLGGVKKWHRNIGNAVPPLGEGIG
Dnmt2	RYFTPKELIANLQGFP	EKTTVKQRYRLIGNSLNVHVAKLL
M.Spr	RRLTPLECFRLQAFD	AGISNSOLYKQTGNSITVTLESIF
DNMT3A	DILWCTEMERVFGFP	SNMSRLARQRLGRSWSVPVIRHLF
Dnmt3a	DILWCTEMERVFGFP	SNMSRLARQRLGRSWSVPVIRHLF
DNMT3B	DVLWCTELERIFGFP	SNMGRGARQKLLGRSWSVPVIRHLF
Dnmt3b	DVLWCTELERIFGFP	SNMGRGARQKLLGRSWSVPVIRHLF
Zmt3	DHIWITTELEKIFGFP	KSMGRPORQRVLGKSWSPVIRHLL
consensus	-----E-R---GFP	-----R-G---P-----

FIG. 5A

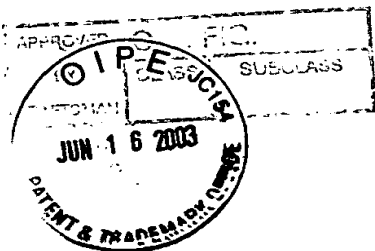
[illegible]

FIG. 5B

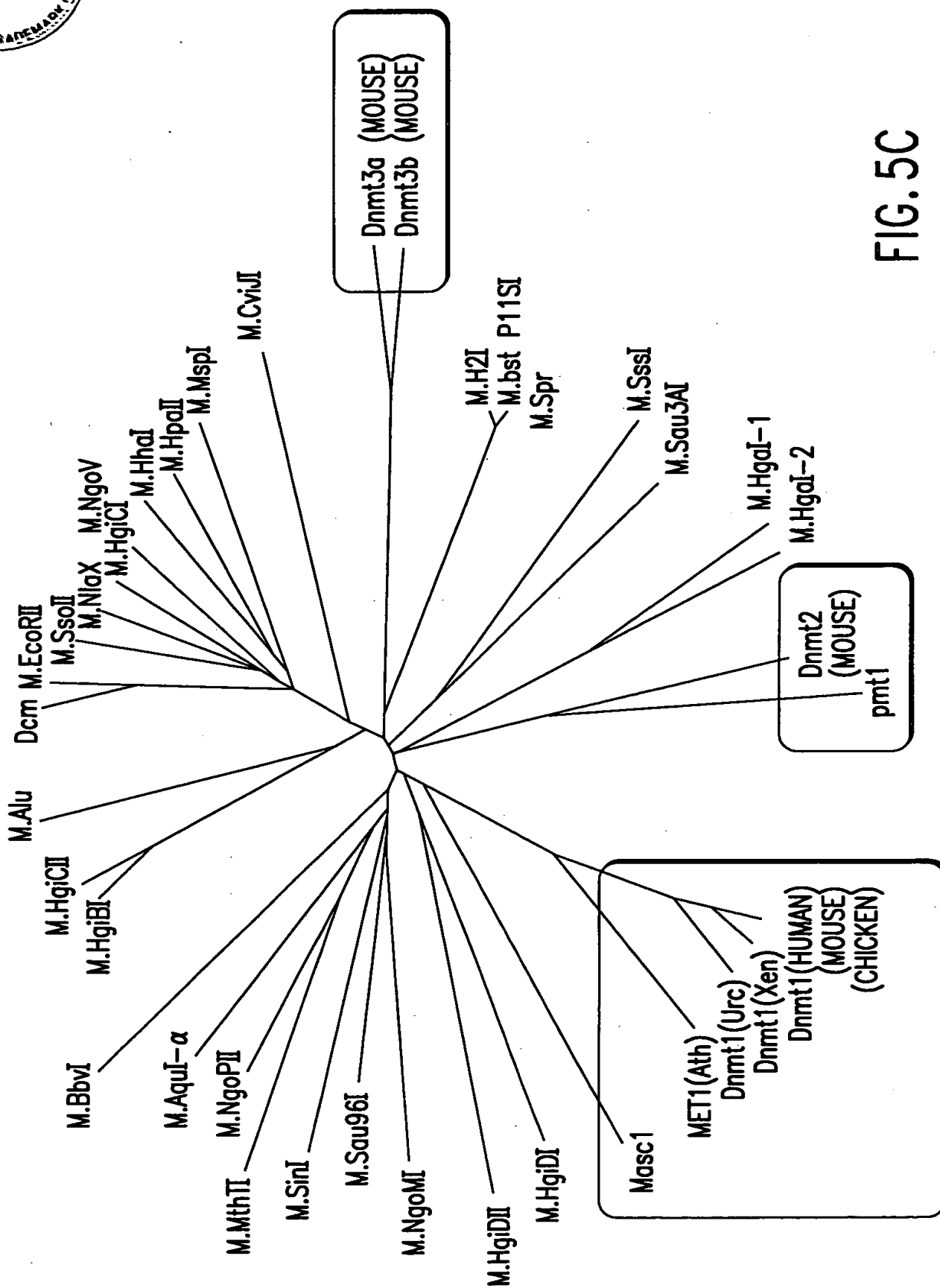
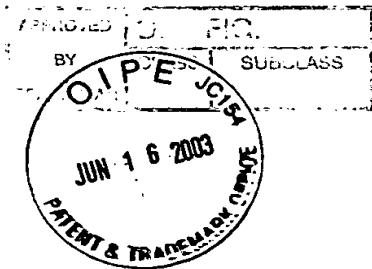


FIG. 5C

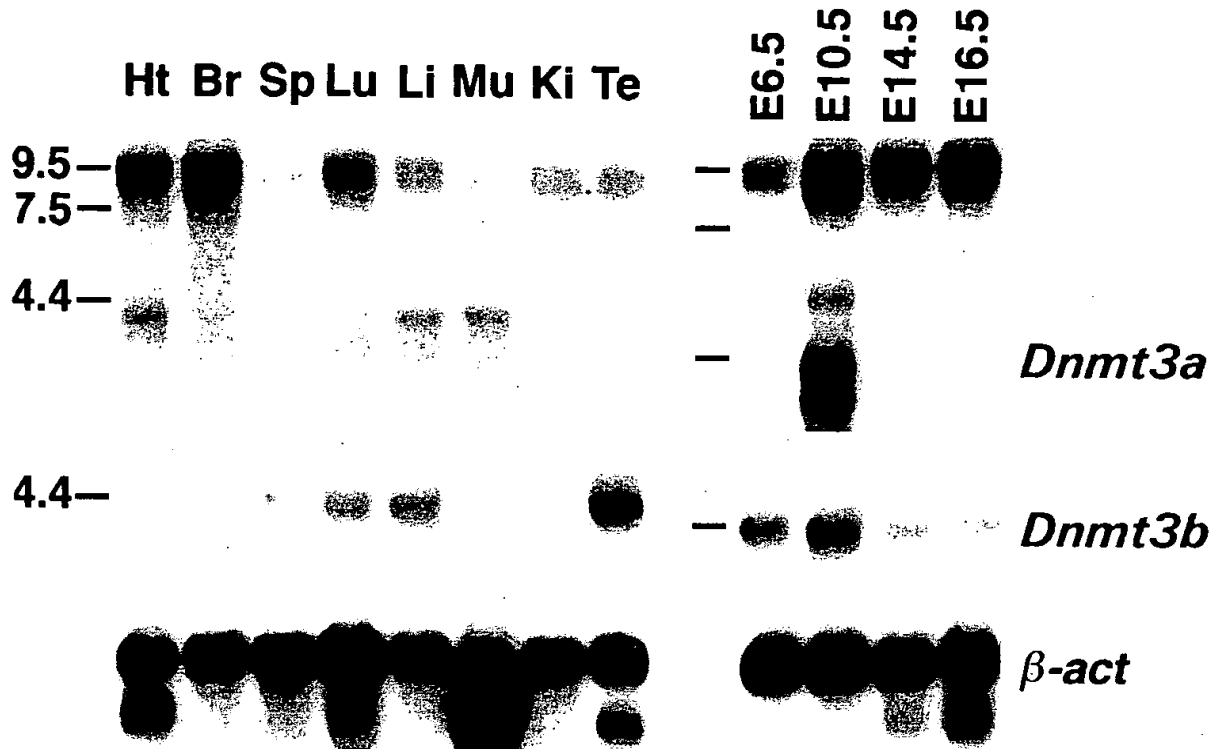


FIG.6A

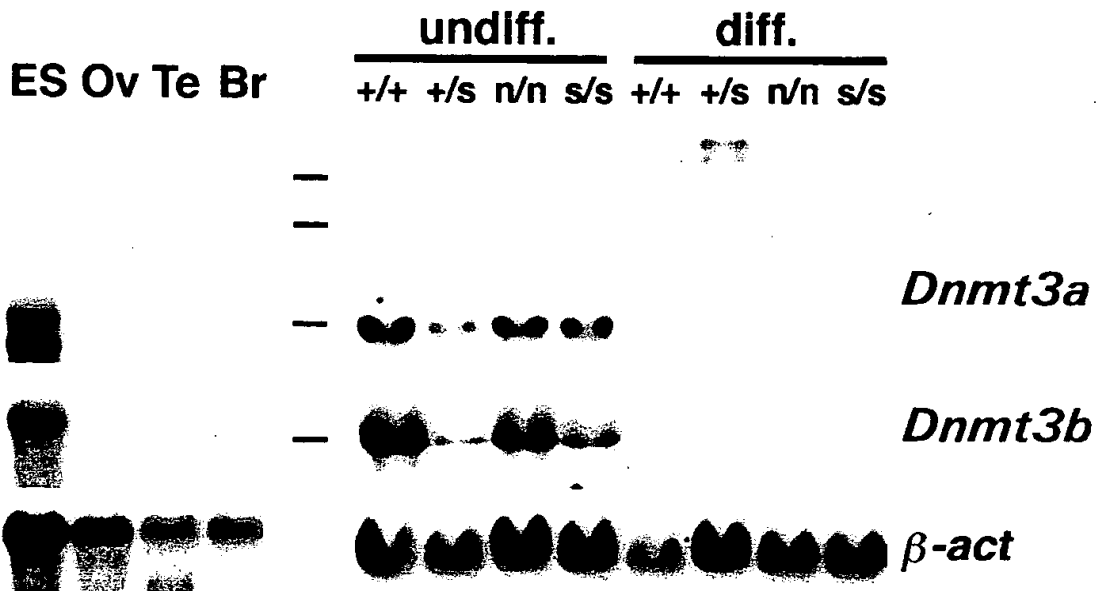


FIG.6B

FIG.6C

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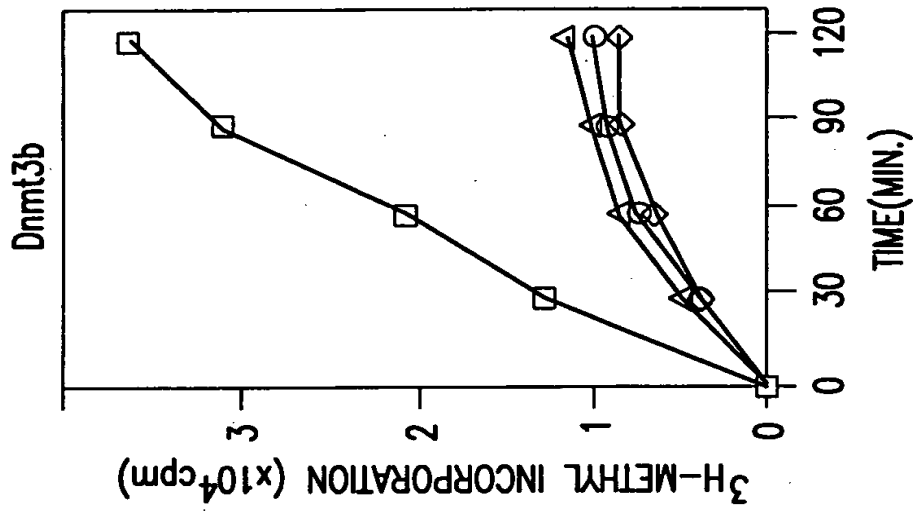
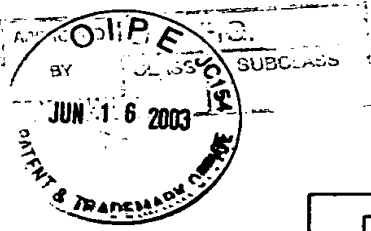


FIG. 7C

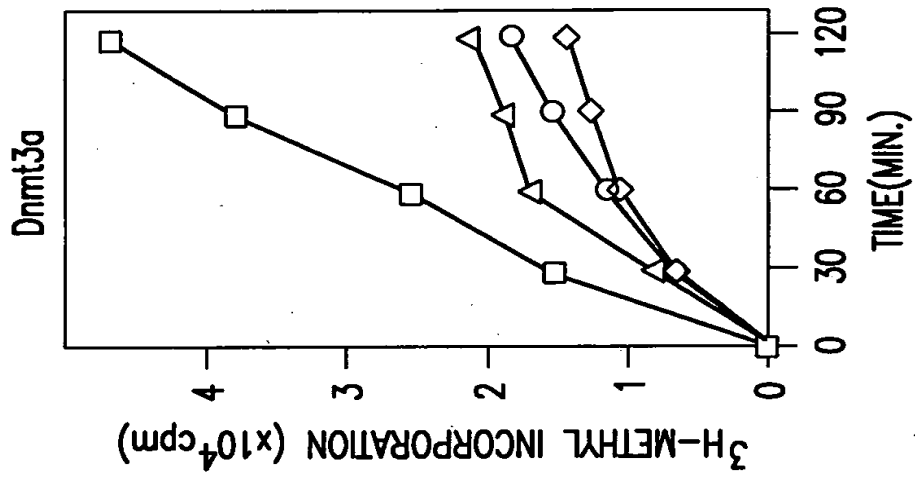


FIG. 7B

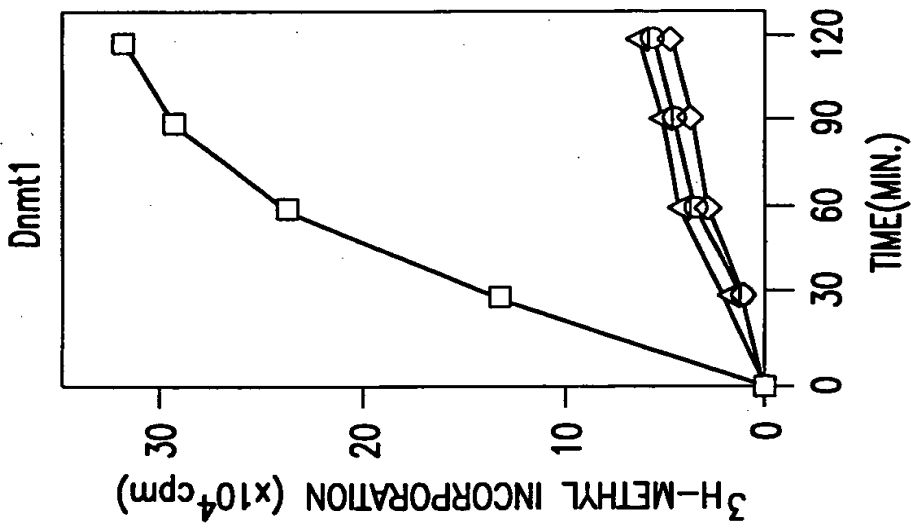


FIG. 7A

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FIG. 7D

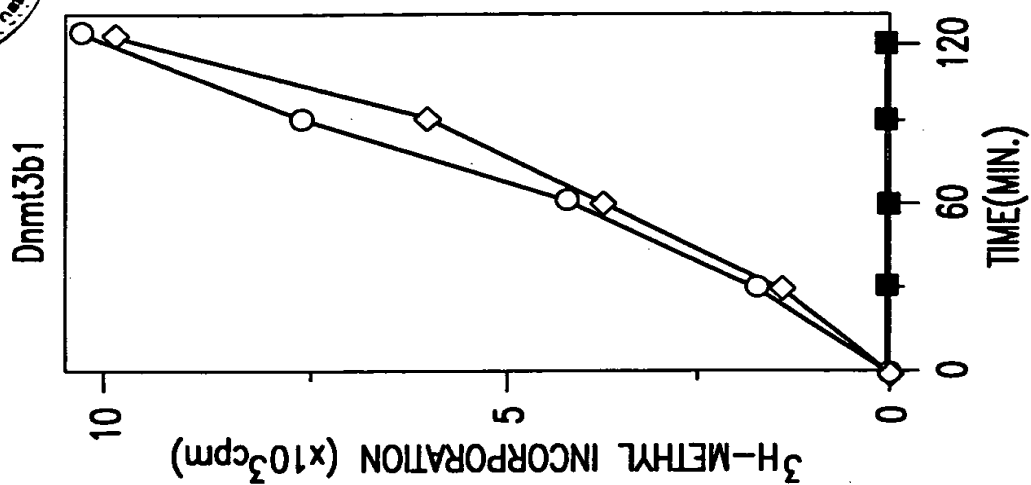


FIG. 8C

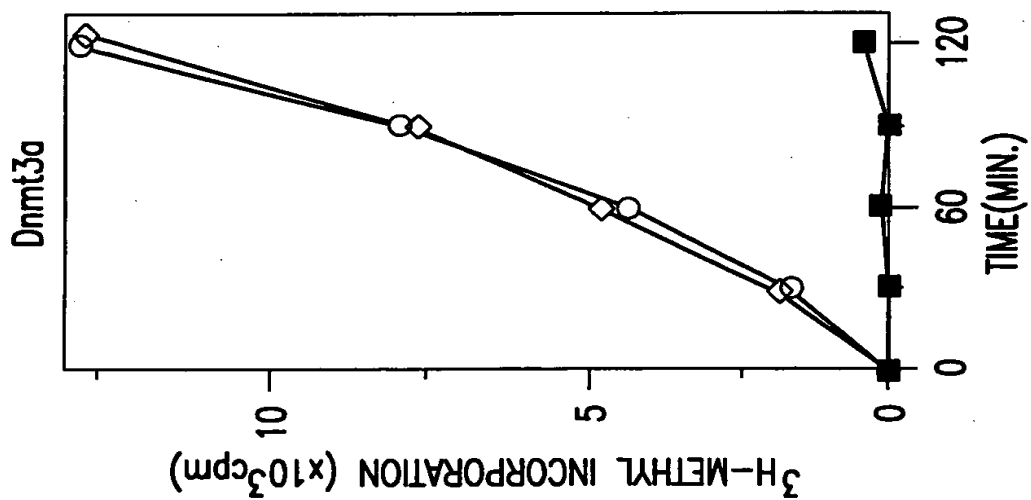


FIG. 8B

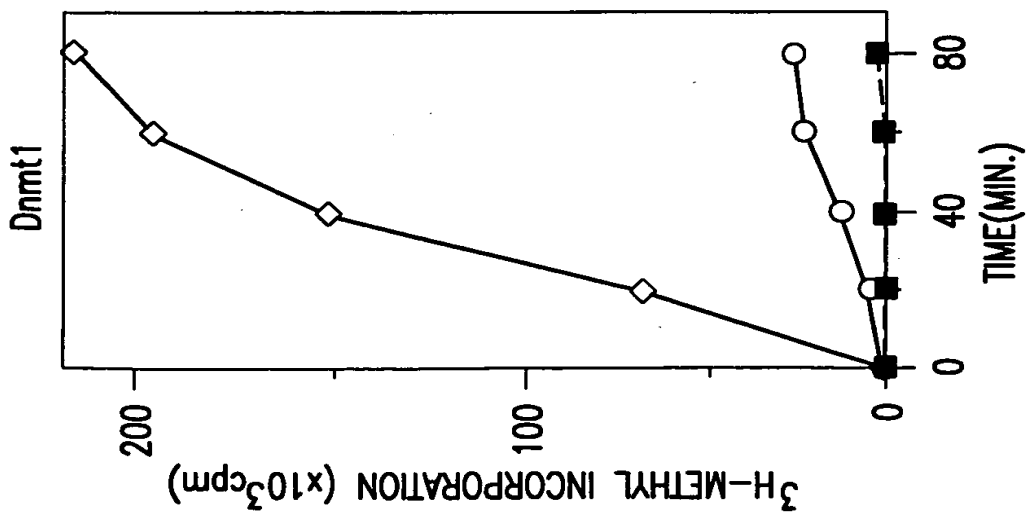


FIG. 8A

APPROVED	C. FIG.	
BY	CLASS	SUBCLASS
DATE/TIME		

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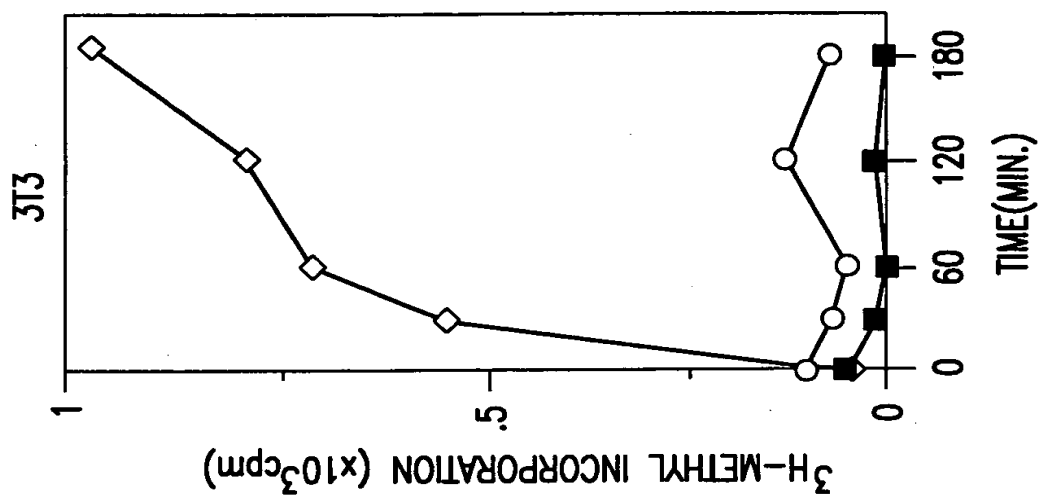


FIG. 8E

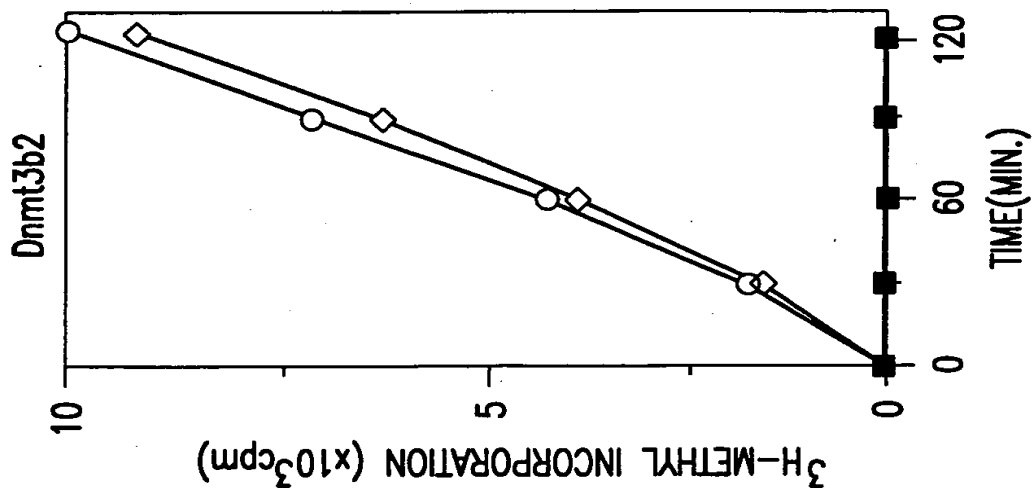
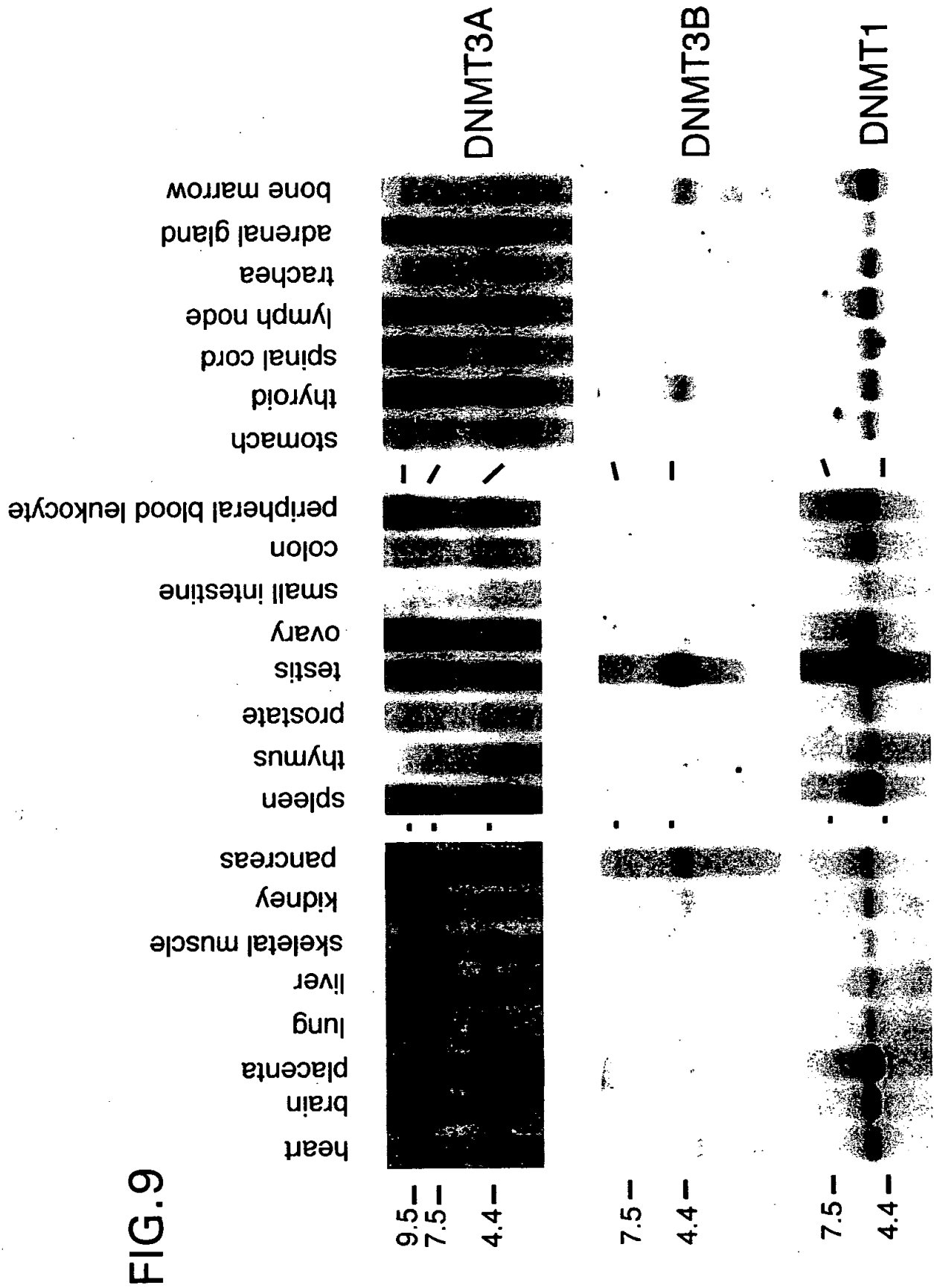


FIG. 8D

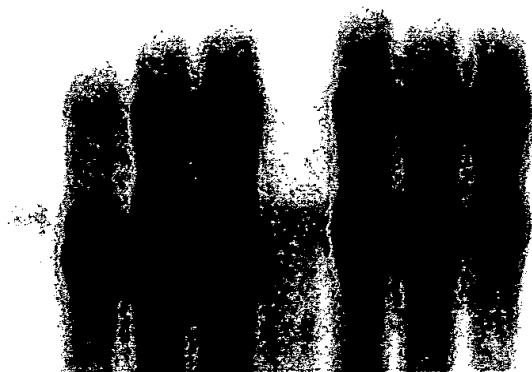


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promyelocytic leukemia HL-60
 Hela cell S3
 chronic myelogenous leukemia K-562
 lymphoblastic leukemia MOLT-4
 Burkitt's lymphoma Raji
 colorectal adenocarcinoma SW480
 lung carcinoma A549
 melanoma G361

9.5 —
 7.5 —
 4.4 —



DNMT3A

7.5 —
 4.4 —



DNMT3B

7.5 —
 4.4 —



DNMT1

2.4 —
 1.4 —



b-Actin

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